

FIG. 1

	A	G	G	C	T	A	G	C	T
GATGGGACTCANGTTCA	1	2	3	4	5	6	7	8	9
GATGGGCTCGNGTTCA	10	11	12	13	14	15	16	17	18
GATGGGCTCCNGTTCA	19	20	21	22	23	24	25	26	27
GATGGGCTCTNGTTCA	28	29	30	31	32	33	34	35	36
GATGGGCTCANGTTCA	37	38	39	40	41	42	43	44	45
GATGGGCTCCNGTTCA	46	47	48	49	50	51	52	53	54
GATGGGCTCTNGTTCA	55	56	57	58	59	60	61	62	63
GATGGGCTCANGTTCA	64	65	66	67	68	69	70	71	72

FIG. 2

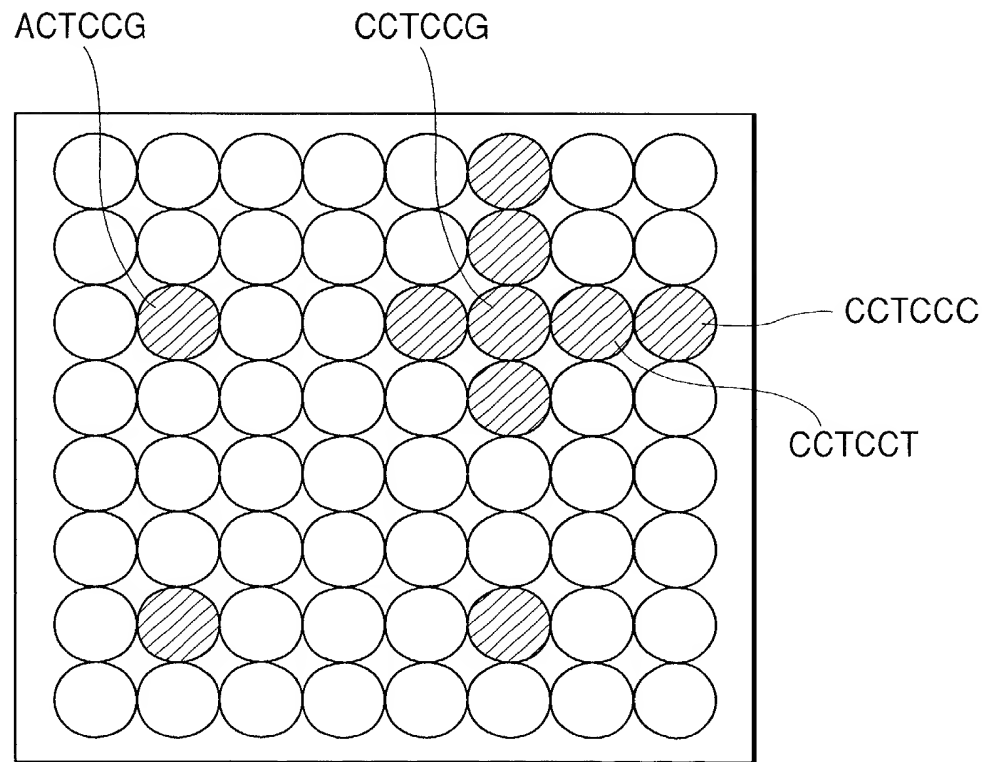
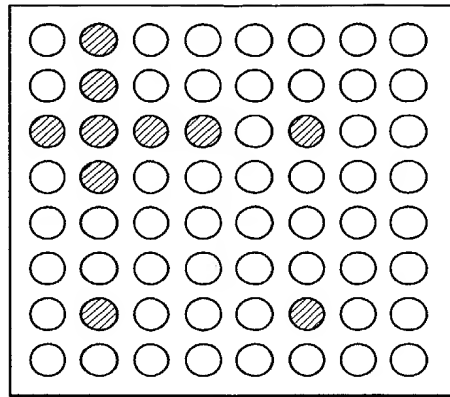
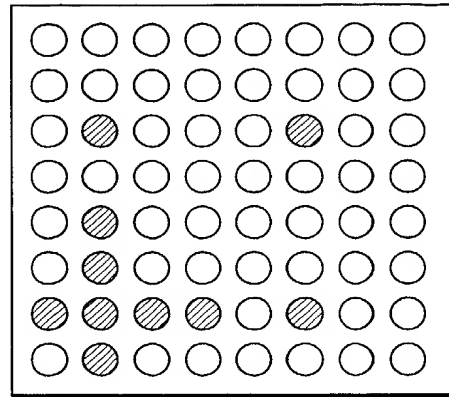
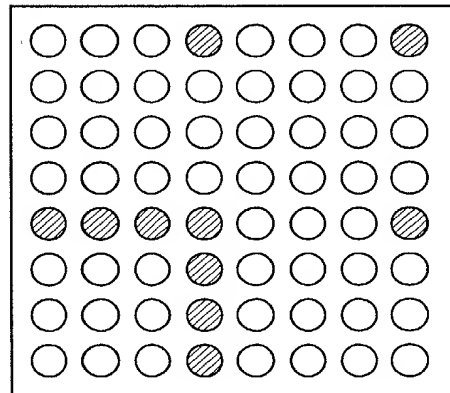
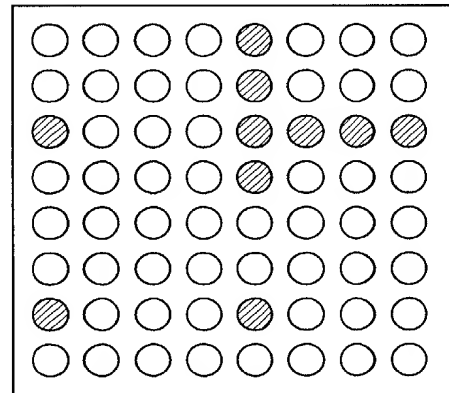
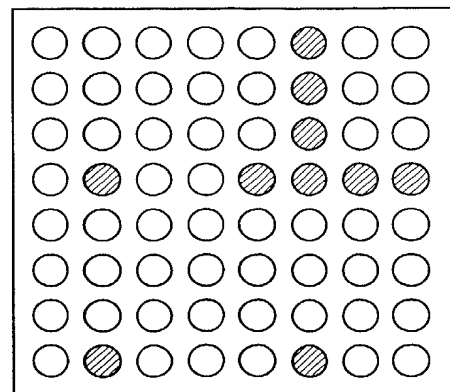
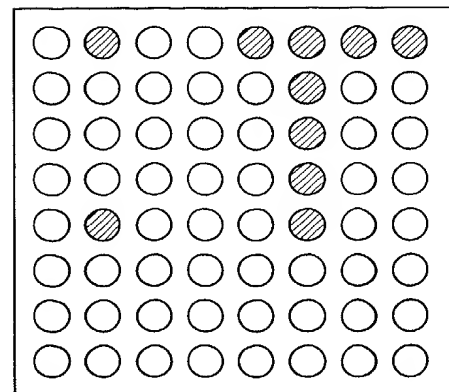


FIG. 3

ATGAACCGGAGTCCCATC

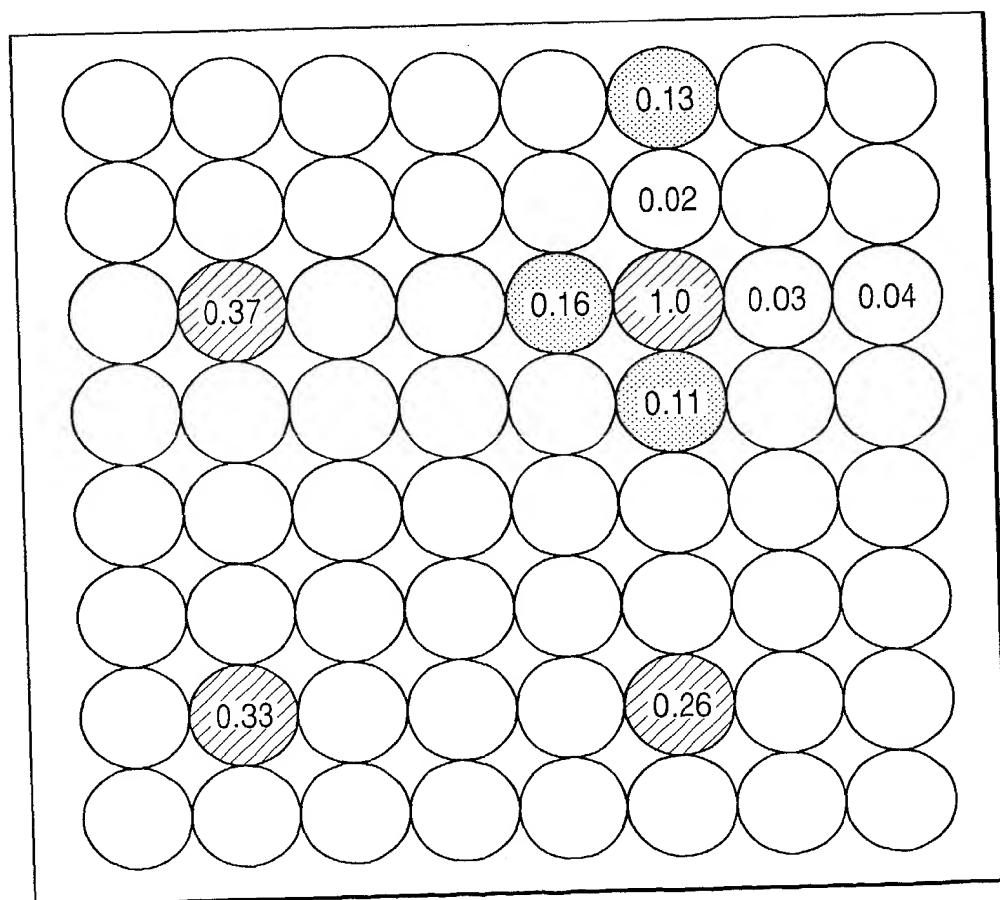
ATGAACCGGAGCCCCATCATGAACATGAGCCCCATCATGAACTGGAGGCCCATC

ATGAACCAGAGGCCCATC

ATGAACCTGAGGCCCATC

TGTGGGAGGAGG

FIG. 4



T07600 9534660

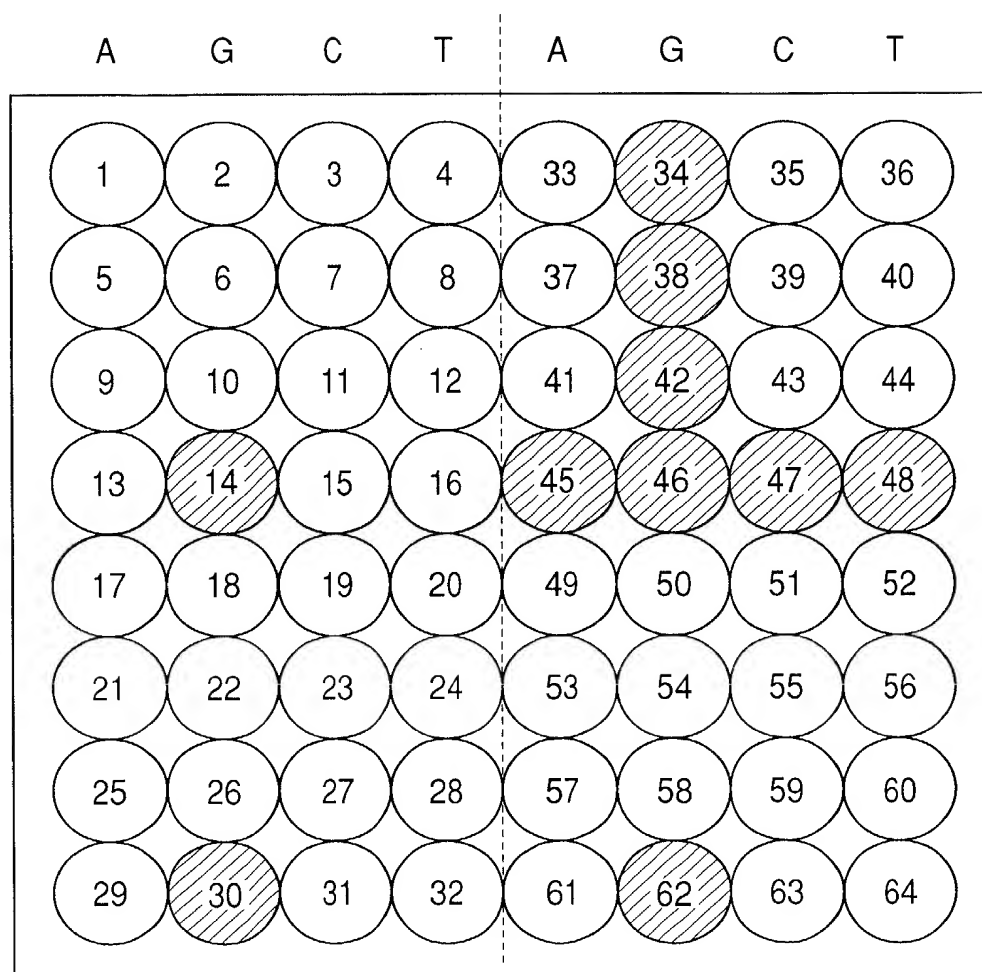
FIG. 5

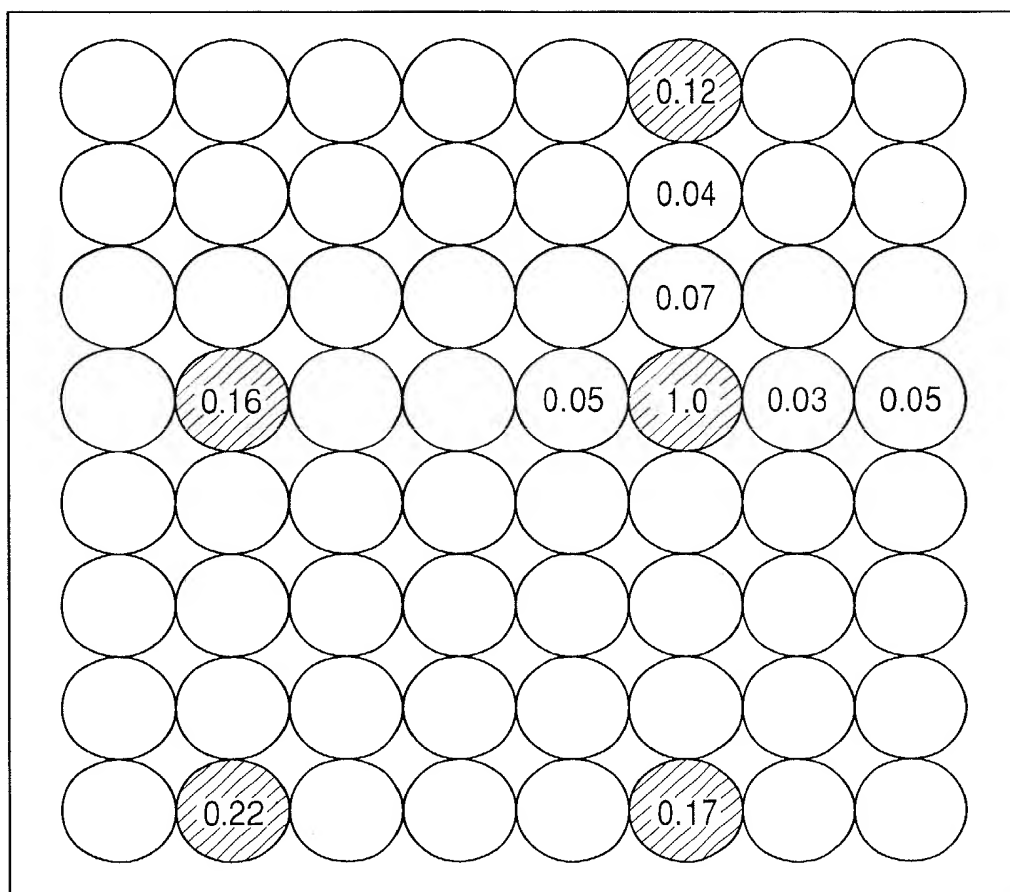
FIG. 6

FIG. 7